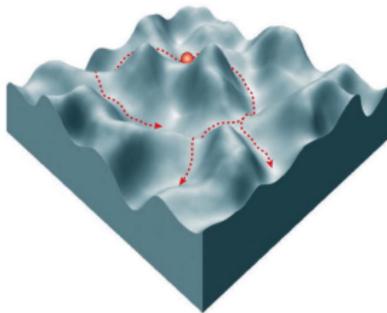


# Taming Asynchrony for Attractor Detection in Large Boolean Networks



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## Biological background & motivations

# Cell potency

Cell potency is a cell's ability to differentiate into other cell types.

- Embryonic stem cells: **pluripotent** – have the potential to differentiate into any of the three germ layers: endoderm, mesoderm, or ectoderm.
- Stem cells derived from adult tissues: **multipotent** – maintain a limited, tissue-specific, regenerative potential

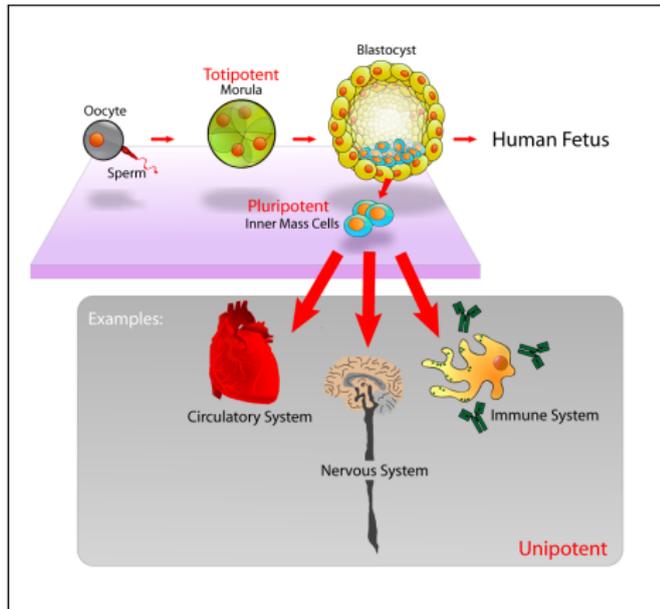
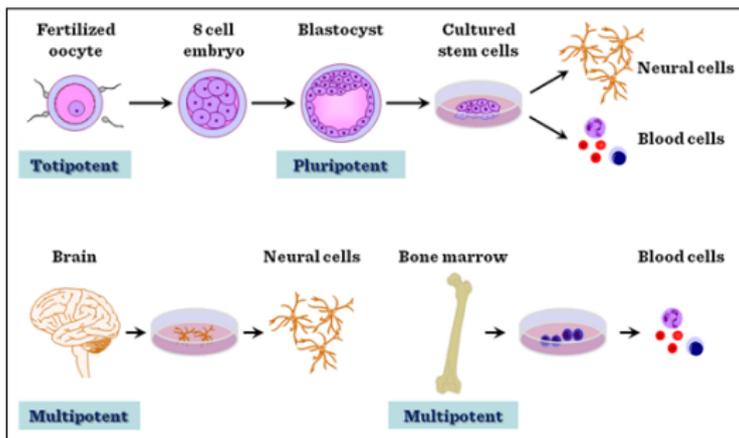


Image: Wikipedia

# Cellular reprogramming

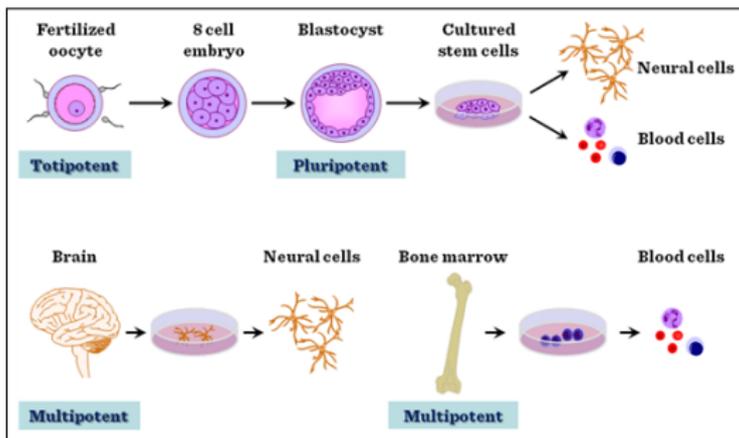
- Reprogramming of differentiated adult cell to embryonic-like pluripotent state
- Reprogramming to other adult cell types **without intermediate reversion** to a pluripotent state



D. Bartis and J. Pongrácz. Three dimensional tissue cultures and tissue engineering. University of Pécs, 2011.

# Cellular reprogramming

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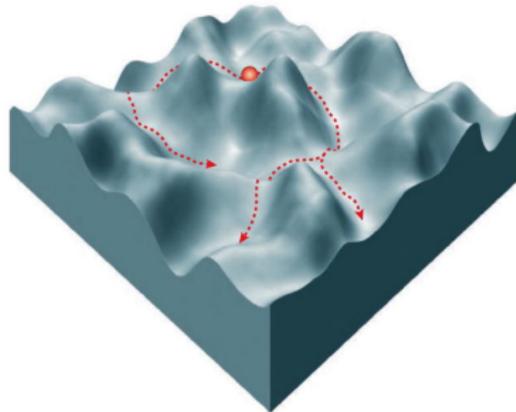
D. Bartis and J. Pongrácz. Three dimensional tissue cultures and tissue engineering. University of Pécs, 2011.

Disease modelling, drug design, organ synthesis, tissue repair, etc.

# Conrad Waddington's epigenetic landscape

- A visual metaphor for the embryonic development and cellular commitment
- Development viewed as a ball rolling down a sloping landscape containing multiple 'hills' and 'valleys'
- Cells take different paths down this landscape and so adopt different fates
- Hills act as barriers by separating the landscape into valleys (**cell types**)
- **Differentiation is not terminal:** epigenetic barriers that can be overcome given sufficient perturbations

Epigenotype – “complex network of processes and causal mechanisms by which the genes of the genotype bring about phenotypic effects”



Phenotype = genotype + epigenotype + environment



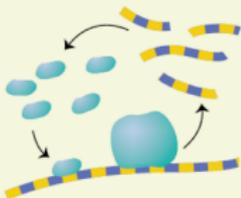
The idea that **cell types** may be related to 'balanced states' of an underlying regulatory system bears **a striking resemblance** to the modern mathematical notion of **attractors of dynamical systems**.

Low

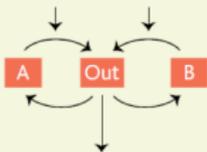
System complexity

High

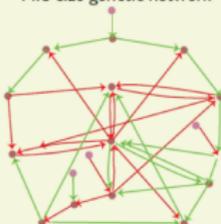
Single gene



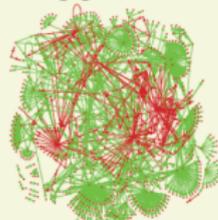
Small genetic circuit



Mid-size genetic network



Large genetic network

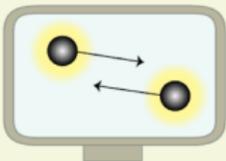


Great detail

Computer modeling

Less detail

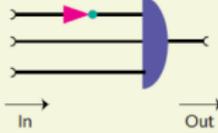
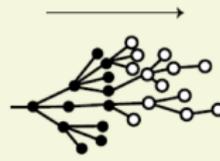
Stochastic molecular simulations



Differential equations

$$\frac{d[RNA]}{dt} = \frac{\alpha k^h}{k^h + [Protein(t-\tau)]}$$

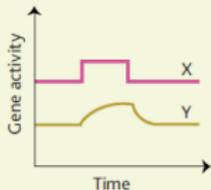
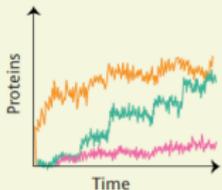
$$\frac{d[Protein]}{dt} = \beta[RNA(t)]$$

Discrete dynamics  
(connected switches)Pseudodynamics  
(flow across a network)

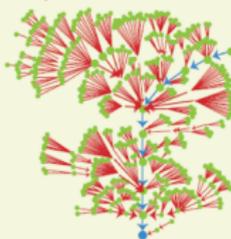
Single gene dynamics

Information

System dynamics



Flow pattern of network states



Functional modules

S. Bornholdt. **Less Is More in Modeling Large Genetic Networks.**

Science, 310(5747):449-451, 2005

## Boolean networks (BNs)

# Boolean networks

Boolean networks are a class of discrete dynamical systems that can be characterised by interactions over a set of Boolean variables.

- First introduced by Stuart Kauffman in 1969 as a simple model class for studying dynamical properties of gene regulatory networks.
- Assumption: genes can be in two possible states of activity (e.g., ON or OFF) at any given point in time, and that they act on each other by means of Boolean functions.

## Boolean networks

- A **Boolean network**  $G(V, \mathbf{f})$  is defined as a set of binary valued variables (also referred to as nodes or genes),  $V = \{x_1, x_2, \dots, x_n\}$  and a vector of Boolean functions  $\mathbf{f} = (f_1, f_2, \dots, f_n)$ .
- At each time point  $t$ , the **state** of the network is defined by the vector  $\mathbf{x}(t) = (x_1(t), x_2(t), \dots, x_n(t))$ , where  $x_i(t)$  is the value of variable  $x_i$  at time  $t$ , i.e.,  $x_i(t) \in \{0, 1\}$ .
- For each variable  $x_i$ , there exists a **predictor set**  $\{x_{i_1}, x_{i_2}, \dots, x_{i_{k(i)}}\}$ , and a Boolean **predictor function** (or simply **predictor**)  $f_i$  being the  $i$ -th element of  $\mathbf{f}$  that determines the value of  $x_i$  at the next time point, i.e.,

$$x_i(t+1) = f_i(x_{i_1}(t), x_{i_2}(t), \dots, x_{i_{k(i)}}(t)),$$

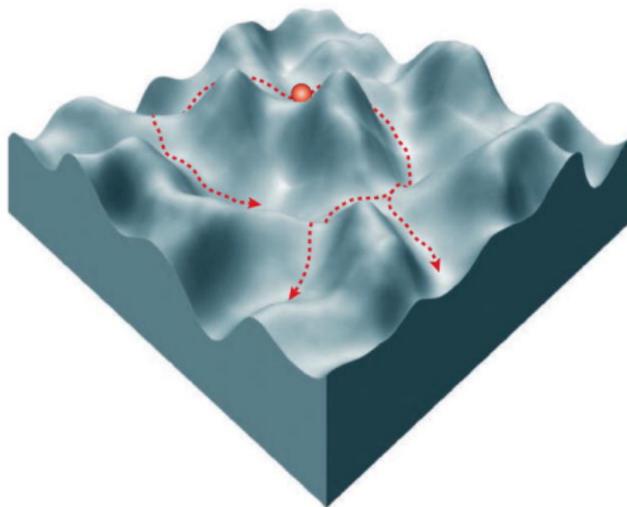
where  $1 \leq i_1 < i_2 < \dots < i_{k(i)} \leq n$ .

## Synchronous Boolean networks: Attractors

$$\mathbf{x}(t + 1) = \mathbf{f}(\mathbf{x}(t))$$

- Given an initial state, within a finite number of steps, the BN will transition into: a fixed state, called **singleton attractor**, or a set of states through which it will repeatedly cycle forever, referred to as **cyclic attractor**.
- The **attractor structure** of a BN is determined by the particular combination of singleton and cyclic attractors, and by the cycle lengths of the cyclic attractors. The attractors of a BN characterise its **long-run behaviour**.
- The states that lead into an attractor constitute its **basin of attraction**. The basins form a partition of the state space.

# Attractor detection for large asynchronous BNs



# BDD-based attractor detection for asynchronous BNs

The **asynchronous updating scheme**:  $T(x(t), x(t + 1)) =$

$$\left( x_i(t + 1) \leftrightarrow f_i(x_{i_1}(t), x_{i_2}(t), \dots, x_{i_{k_i}}(t)) \right) \wedge_{j \neq i} (x_j(t + 1) \leftrightarrow x_j(t))$$

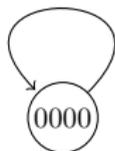
It states that node  $v_i$  is updated by its Boolean function and other nodes are kept unchanged.

## BDD-based attractor detection for asynchronous BNs

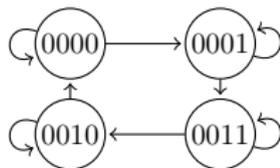
[**Attractor of a BN**] An **attractor** of a BN is a set of states satisfying that any state in this set can be reached from any other state in this set and no state in this set can reach any other state that is not in this set.

[**Attractor system**] An attractor together with its state transition relations is referred to as an **attractor system**.

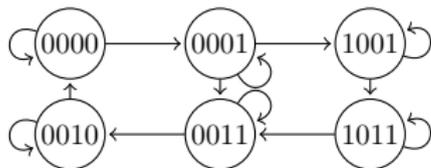
Three types of attractor systems for asynchronous BNs (BSCCs):



(a) A selfloop



(b) A simple loop



(c) A complex loop

# BDD-based attractor detection for asynchronous BNs

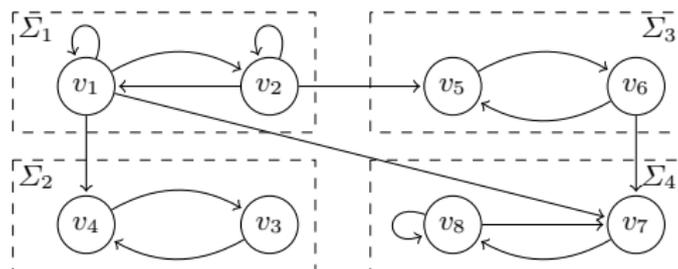
[Block] Given a BN  $G(V, \mathbf{f})$  with  $V = \{v_1, v_2, \dots, v_n\}$  and  $\mathbf{f} = \{f_1, f_2, \dots, f_n\}$ , a **block**  $B(V^B, \mathbf{f}^B)$  is a subset of the network, where  $V^B \subseteq V$ . For any node  $v_i \in V^B$ , if  $B$  contains all the parent nodes of  $v_i$ , its Boolean function in  $B$  remains the same as in  $G$ , i.e.,  $f_i$ ; otherwise, the Boolean function is undetermined, meaning that additional information is required to determine the value of  $v_i$  in  $B$ .

We call the nodes with undetermined Boolean functions as **undetermined nodes**.

We refer to a block as an **elementary block** if it contains no undetermined nodes.

# BDD-based attractor detection for asynchronous BNs

Our decomposition method:



- An elementary block contains no undetermined nodes.
- Given a BN  $G$ , let  $B$  be an elementary block in  $G$ .  $B$  preserves the attractors of  $G$ .

## BDD-based attractor detection for asynchronous BNs

[Projection map, Compressed state, Mirror states] For a BN  $G$  and its block  $B$ , where the set of nodes in  $B$  is  $V^B = \{v_1, v_2, \dots, v_m\}$  and the set of nodes in  $G$  is  $V = \{v_1, v_2, \dots, v_m, v_{m+1}, \dots, v_n\}$ , the **projection map**  $\pi_B : X \rightarrow X^B$  is given by  $\mathbf{x} = (x_1, x_2, \dots, x_m, x_{m+1}, \dots, x_n) \mapsto \pi_B(\mathbf{x}) = (x_1, x_2, \dots, x_m)$ .

For any set of states  $S \subseteq X$ , we define  $\pi_B(S) = \{\pi_B(\mathbf{x}) : \mathbf{x} \in S\}$ . The projected state  $\pi_B(\mathbf{x})$  is called a **compressed state** of  $\mathbf{x}$ . For any state  $\mathbf{x}^B \in X^B$ , we define its set of **mirror states** in  $G$  as  $\mathcal{M}_G(\mathbf{x}^B) = \{\mathbf{x} \mid \pi_B(\mathbf{x}) = \mathbf{x}^B\}$ . For any set of states  $S^B \subseteq X^B$ , its set of mirror states is  $\mathcal{M}_G(S^B) = \{\mathbf{x} \mid \pi_B(\mathbf{x}) \in S^B\}$ .

## BDD-based attractor detection for asynchronous BNs

[**Preservation of attractors**] Given a BN  $G$  and an elementary block  $B$  in  $G$ , let  $\mathcal{A} = \{A_1, A_2, \dots, A_m\}$  be the set of attractors of  $G$  and  $\mathcal{A}^B = \{A_1^B, A_2^B, \dots, A_{m'}^B\}$  be the set of attractors of  $B$ . We say that  $B$  **preserves the attractors** of  $G$  if for any  $k \in [1, m]$ , there is an attractor  $A_{k'}^B \in \mathcal{A}^B$  such that  $\pi_B(A_k) \subseteq A_{k'}^B$ .

# BDD-based attractor detection for asynchronous BNs

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## Lemma

*Given a BN  $G$  and an elementary block  $B$  in  $G$ , let  $\Phi$  be the set of attractor states of  $G$  and  $\Phi^B$  be the set of attractor states of  $B$ . If  $B$  preserves the attractors of  $G$ , then  $\Phi \subseteq \mathcal{M}_G(\Phi^B)$ .*

## Theorem

*Given a BN  $G$ , let  $B$  be an elementary block in  $G$ .  $B$  preserves the attractors of  $G$ .*

# BDD-based attractor detection for asynchronous BNs

## [Crossability, Cross operations]

- $B_i$  – a non-elementary block,  
 $V^{B_i} = \{v_{p_1}, v_{p_2}, \dots, v_{p_s}, v_{q_1}, v_{q_2}, \dots, v_{q_t}\}$
- $B_j$  –  $B_i$ 's parent block,  
 $V^{B_j} = \{v_{q_1}, v_{q_2}, \dots, v_{q_t}, v_{r_1}, v_{r_2}, \dots, v_{r_u}\}$
- $\mathbf{x}^{B_i} = (x_1, x_2, \dots, x_s, y_1^i, y_2^i, \dots, y_t^i)$  a state of  $B_i$
- $\mathbf{x}^{B_j} = (y_1^j, y_2^j, \dots, y_t^j, z_1, z_2, \dots, z_u)$  a state of  $B_j$
- **Crossable:**  $\mathbf{x}^{B_i} \mathcal{C} \mathbf{x}^{B_j}$ , if the values of their common nodes are the same, i.e.,  $y_k^i = y_k^j$  for all  $k \in [1, t]$ .
- **The cross operation:**  
 $\Pi(\mathbf{x}^{B_i}, \mathbf{x}^{B_j}) = (x_1, x_2, \dots, x_s, y_1^i, y_2^i, \dots, y_t^i, z_1, z_2, \dots, z_u)$

# BDD-based attractor detection for asynchronous BNs

## [Crossability, Cross operations]

We say a set of states  $S^{B_i} \subseteq X^{B_i}$  and a set of states  $S^{B_j} \subseteq X^{B_j}$  are crossable, denoted as  $S^{B_i} \mathcal{C} S^{B_j}$ , if at least one of the sets is empty or the following two conditions hold: 1) for any state  $\mathbf{x}^{B_i} \in S^{B_i}$ , there always exists a state  $\mathbf{x}^{B_j} \in S^{B_j}$  such that  $\mathbf{x}^{B_i}$  and  $\mathbf{x}^{B_j}$  are crossable; 2) vice versa.

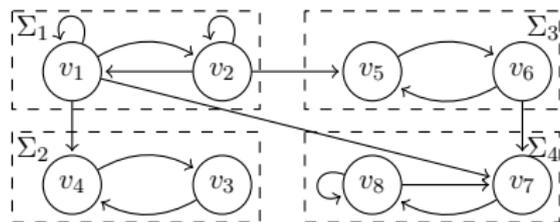
$$\Pi(S^{B_i}, S^{B_j}) = \{\Pi(\mathbf{x}^{B_i}, \mathbf{x}^{B_j}) \mid \mathbf{x}^{B_i} \in S^{B_i}, \mathbf{x}^{B_j} \in S^{B_j} \text{ and } \mathbf{x}^{B_i} \mathcal{C} \mathbf{x}^{B_j}\}.$$

Let  $\mathcal{S}^{B_i} = \{S^{B_i} \mid S^{B_i} \subseteq X^{B_i}\}$  be a family of states sets in  $B_i$  and  $\mathcal{S}^{B_j} = \{S^{B_j} \mid S^{B_j} \subseteq X^{B_j}\}$  be a family of states sets in  $B_j$ . We say  $\mathcal{S}^{B_i}$  and  $\mathcal{S}^{B_j}$  are crossable, denoted as  $\mathcal{S}^{B_i} \mathcal{C} \mathcal{S}^{B_j}$  if for any states set  $S^{B_i} \in \mathcal{S}^{B_i}$ , there always exists a states set  $S^{B_j} \in \mathcal{S}^{B_j}$  such that  $S^{B_i}$  and  $S^{B_j}$  are crossable; 2) vice versa.

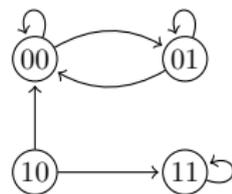
$$\Pi(\mathcal{S}^{B_i}, \mathcal{S}^{B_j}) = \{\Pi(S_i, S_j) \mid S_i \in \mathcal{S}^{B_i}, S_j \in \mathcal{S}^{B_j} \text{ and } S_i \mathcal{C} S_j\}.$$

# BDD-based attractor detection for asynchronous BNs

[Realisation of a block]



(a) SCC decomposition of a BN.



(b) Transition graph of  $B_1$ .

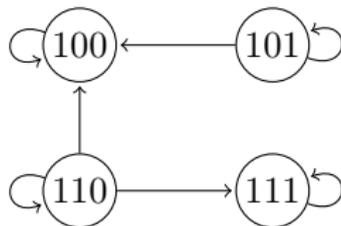
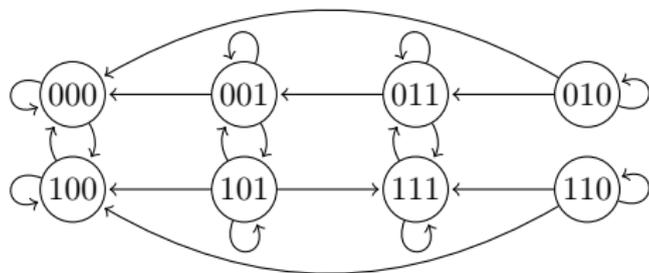


Figure: Transition graphs of two realisations of  $B_3$ .

# BDD-based attractor detection for asynchronous BNs

## Main theorem

[Crossability of attractors] Let  $B_i$  be a single and elementary parent block of a non-elementary block  $B_j$  in a BN  $G$ . Let  $A^{B_i}$  be an attractor of  $B_i$  and let  $A^{B_j}$  be an attractor in the realisation of  $B_j$  with respect to  $A^{B_i}$ . Then  $A^{B_i} \subset A^{B_j}$ .

[Attractor recovery] Given a BN  $G$  with  $B_i$  and  $B_j$  being its two blocks, let  $\mathcal{A}^{B_i}$  and  $\mathcal{A}^{B_j}$  be the set of attractors for  $B_i$  and  $B_j$ , respectively.

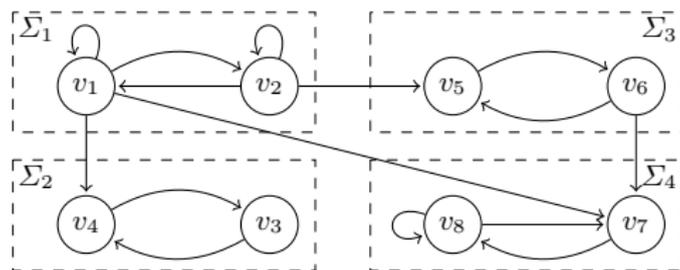
Let  $B_{i,j}$  be the block got by merging the nodes in  $B_i$  and  $B_j$ . If  $B_i$  and  $B_j$  are both elementary blocks or  $B_i$  is an elementary and single parent block of  $B_j$ , then  $\mathcal{A}^{B_i} \subset \mathcal{A}^{B_j}$  and  $\Pi(\mathcal{A}^{B_i}, \mathcal{A}^{B_j})$  is the set of attractors of  $B_{i,j}$ .

# BDD-based attractor detection for asynchronous BNs

## Corollary

*Given a BN  $G$  with  $B_i$ ,  $B_j$ , and  $B_k$  being its three blocks, let  $\mathcal{A}^{B_i}$ ,  $\mathcal{A}^{B_j}$ , and  $\mathcal{A}^{B_k}$  be the sets of attractors for blocks  $B_i$ ,  $B_j$ , and  $B_k$ , respectively. If the three blocks are all elementary blocks or  $B_i$  is an elementary block and it is the only parent block of  $B_j$  and  $B_k$ , it holds that  $\Pi(\Pi(\mathcal{A}^{B_i}, \mathcal{A}^{B_j}), \mathcal{A}^{B_k}) = \Pi(\Pi(\mathcal{A}^{B_i}, \mathcal{A}^{B_k}), \mathcal{A}^{B_j})$ .*

# BDD-based attractor detection for asynchronous BNs



- Compute the attractors of the elementary block  $B_1$ .
- For  $B_2$ , form realisations of  $B_2$  with the attractors of  $B_1$ , then compute attractor of each of the realisations.
- Similarly, compute the attractors for  $B_3$ .
- $B_4$  has two parent blocks: merge  $B_1$  and  $B_3$  as its new parent block, and compute its attractors.
- Similarly, compute the attractors for  $B_4$ .
- Recover the attractors of the BN by cross operations.

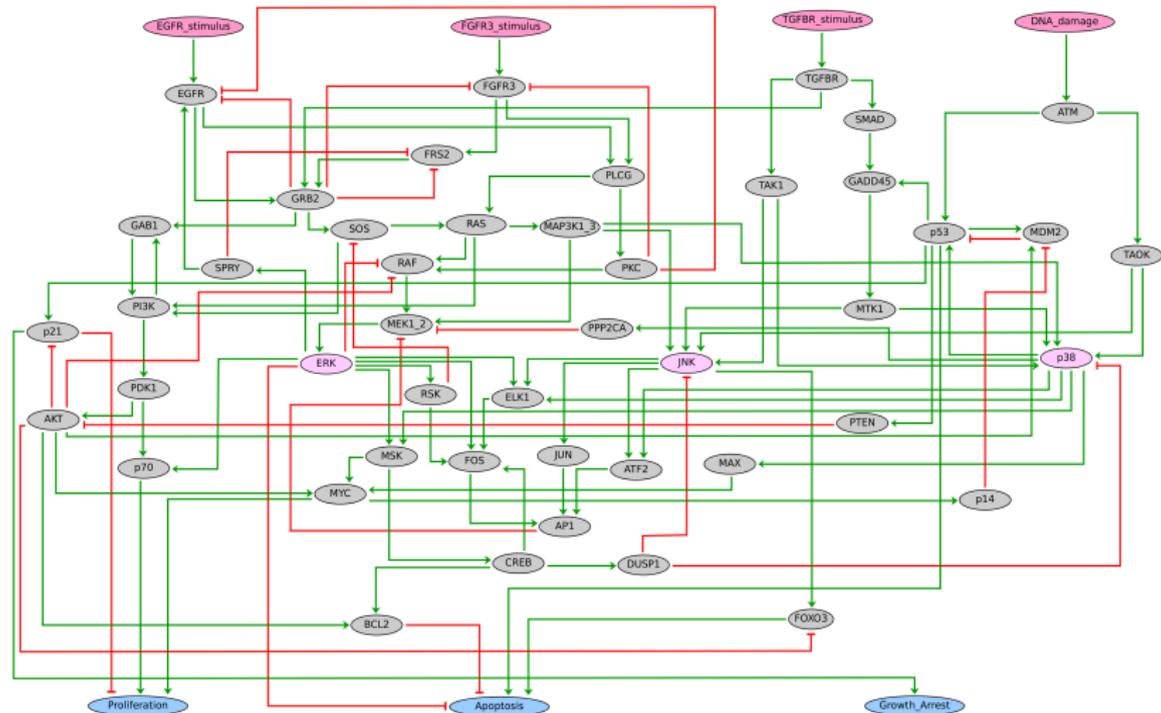
# BDD-based attractor detection for asynchronous BNs

**Evaluation:** MAPK network (53 nodes) & apoptosis (97 nodes)

Networks	# attractors	Time(s)		Speedup	Networks	# attractors	Time(s)		Speedup
		Alg. 1	Alg. 2				Alg. 1	Alg. 2	
MAPK_r3	20	6.070	2.614	2.32	apoptosis	1024	1633.970	103.856	15.73
MAPK_r4	24	11.674	1.949	5.99	apoptosis*	2048	8564.680	218.230	39.25

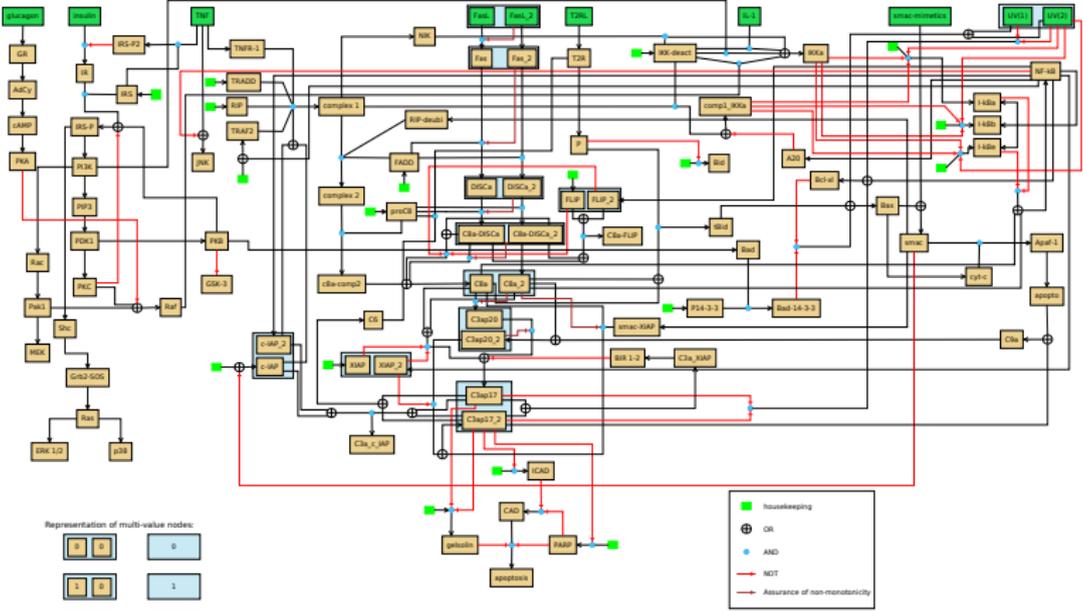
# BDD-based attractor detection for asynchronous BNs

## MAPK network



# BDD-based attractor detection for asynchronous BNs

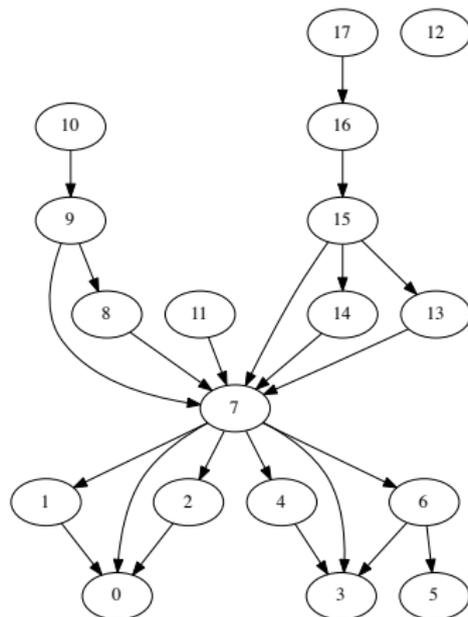
## Apoptosis network



To minimize this option the input of the first subnode should be combined via an AND operator with the negation of the input of the second subnode for each multi-valued node.  
The two subnodes of each of the input nodes Tnfα and CD95 are meant to be in the same sense.

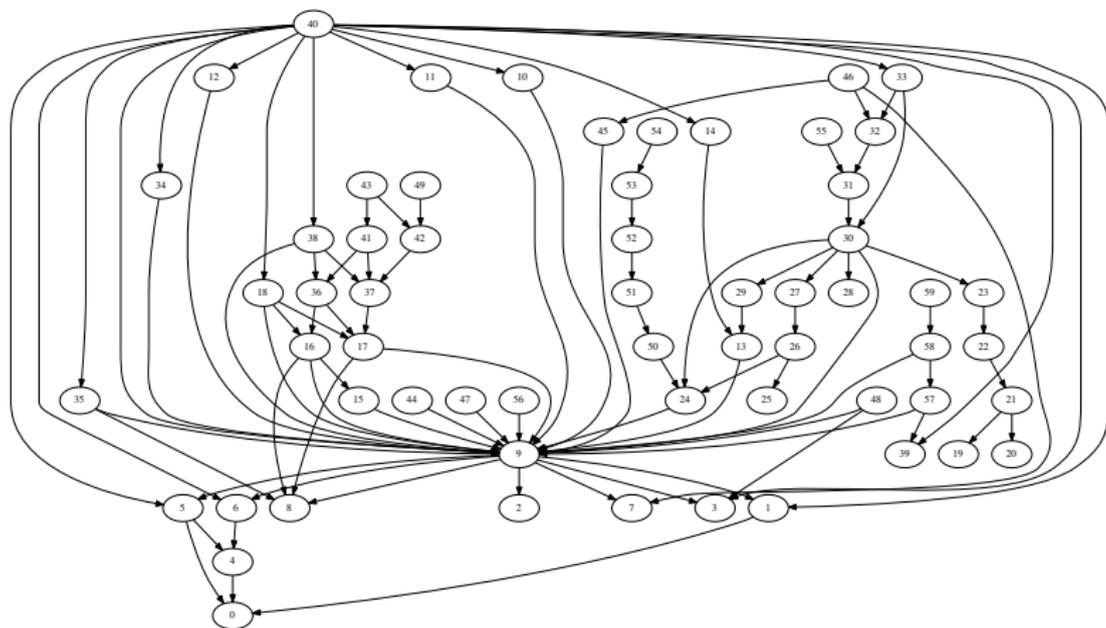
# BDD-based attractor detection for asynchronous BNs

## MAPK network SCC decomposition



# BDD-based attractor detection for asynchronous BNs

## Apoptosis network SCC decomposition



Software tool: ASSA-PBN

## ASSA-PBN

**A**pproximate **S**tead-**S**tate **A**nalyser-PBN

a software tool for modelling, simulation and analysis of  
probabilistic Boolean networks

Freely available at

<http://satoss.uni.lu/software/ASSA-PBN/>

# Future perspectives



## Scalable control of large biological networks!

- **Network controllability** (e.g., minimal interventions driving the system from a certain attractor to a specific target attractor)
- Two running projects: **SEC-PBN** and **AlgoReCell**

# Acknowledgements



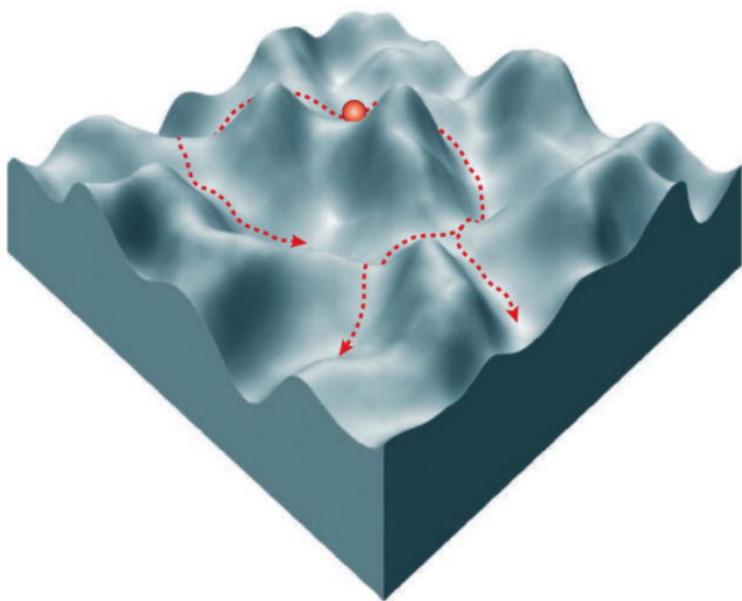
Jun Pang



Hongyang Qu



Qixia Yuan



Thank you for your attention!